

ABSTRACT

The present disclosure provides methods for the diagnosis of metastatic prostate cancer and/or the prediction of the metastatic ability of prostate cancer in prostate biopsy tissue. Metastatic ability of prostate cancer is positively correlated with the level of transcriptional and translational expression of the MUC18 coding sequence in the neoplastic tissue. Methods for the determination of MUC18 protein synthesis include Western blots, ELISA, radioimmunoassay, immunofluorescence, and other immunoassays using MUC18-specific antibody and suitable detection means. Methods for measurement of transcriptional expression of the MUC18 coding sequence include Northern hybridizations and quantitative reverse transcriptase-polymerase chain reaction analyses. Absence of or very low MUC18 expression in the prostate tumor tissue is associated with nonmetastatic cancer, while relatively high levels of MUC18 expression are predictive of prostate cancer which is likely to metastasize or which has already metastasized. The present disclosure provides an improved diagnostic tool to aid the medical community in the choice of appropriate treatment regimens for prostate cancer patients.